

SEQUENCE LISTING

<110> NELSON, DAVID R. <120> A LIVE, AVIRULENT STRAIN OF V. ANGUILLARUM THAT PROTECTS FISH AGAINST INFECTION BY VIRULENT V. ANGUILLARUM AND METHOD FOR MAKING THE SAME <130> 5112 <140> 09/915,706 <141> 2001-07-26 <150> 60/220,733 <151> 2000-07-26 <160> 7 <170> PatentIn Ver. 2.1 <210> 1 <211> 3588 <212> DNA <213> Vibrio anguillarum <220> <221> modified_base <222> (3572) <223> a, t, c, g, other or unknown <400> 1 gtcgacttat tgcattgatg gcgtacatgg tagtgccatc cttcgtttgc taacaagcgt 60 tgtataaaag cttggtcggt ttcatcaagt tgaacacaat actcatgatt tttcccactt 120 ccggaaaggg aaaagtgaaa atagcttttg agatcagcct gttctagcag cttttcaatg 180 atctttttcg tcgttacgtt ttgaaaaatc tgacgactgc gtttgtattg caacaagcta 240 agtggatcca atatctctat ttgataataa aactgctgct tgtctttgct atatcctgtg 300 aattgcagag tgctacatat acctgaaaaa aaacgctttc cagaatctaa ttcgtaagac 360 acacaaacag ctttacctag gtttttggta tcgatctcca tgtttgccgc gatggaaacg 420 gaaaactgac acccgccgga tacgctttcc tctccgatta attgcgtgac aatataactt 480 ttgctatctg aaagcttaat ggtgagggag cgggtttggt gctttaattc gttactgctc 540 atattcaatt aattcactat taaataaaca gttctaaaag gctgtttatt ggatgaatat 600 tcgaaattat cacataataa ttgatgctat tattacttgc tgtattggta tcaactttca 660 tgctctatac atgtaatata tttcgagtta gaccttaatt caaggtaatt tgtctattta 720 attattatct gaataatatg taatcgattg ctttgtggtt atttttatgt ttgtttcatt 780 tttaatgacg gtgagcttgt gcattcatat tttttatgat gacaacatct ttgatgaagt 840 atttaagata ttgttaatgc atgaggggtt tgcgtgtatt ttttatatta aatcataata 900 aaatcaacaa tatatgttat tttgtgtctt tttatagtgt tcttttaaag aggtaggatg 960 acctaaaggt cgcctaaata tggcgtaaat tgccattgct ataattcacc tcaaagatac 1020 actattggca aattgacaaa tatgtcactt cgtatgaaac aatattagta gatgttgttt 1080 ttgctgcaaa aataaaaatt tttctggttg aaataactca aggcctctag cgttttcctt 1140 tatcttaaaa tacaggaaat agcgattgaa gttaattgac acttaagcaa atagtcaacc 1200 taacagagca ggaacctatg cctttgtcaa agcatcaaat tgagcaactt tctaaacctc 1260

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Ser Lys Thr Thr Arg Asp Ile Glu Leu Ile Ser Trp Phe Val Ala Ala

85

90

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- Trp Leu Ala Asp Leu Ser Glu Lys His Trp Asp His Leu Asn Pro Val 115 120 125
- Leu Pro Val Glu Thr Leu Lys Ser Asp Asp Asp Lys Gly Lys Glu Arg 130 135 140
- Glu Gln Ala Asp Ala Lys Val Lys Ala Phe Phe Gln Leu Val Gly Asp
- Ser Glu Glu Ser Ser Ile Leu Tyr Ala Pro Val Leu Gln Leu Pro Leu 165 170 175
- Val Gly Glu Val Thr Phe Phe Asp Phe Gln Ser Ala Glu Arg Lys Gly
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- Arg Phe Ala Ile Gln Phe Lys Met Glu Asn Ala Lys Arg Cys Val Thr 210 215 220
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- Gly Ser Gln Ser Thr Asn Phe Gly Phe Ala Lys Ser Leu Leu Thr Arg 245 250 255
- Val Glu Asn Ala Leu Val His Leu Ser Gly Ile Lys Leu Ala Pro Lys 260 265 270
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- Glu Gly Glu Leu Pro Ser His Met Asp Thr Lys His Ile Glu Arg Ile 290 295 300
- Pro Met Ala Ser Glu Gln Ala Gln Thr Val Ser Gln His Leu His Ala 305 310 315 320
- Gly Asn Leu Ser Glu Leu Gly Asn Leu Asn Asn Met Asn Arg Asp Leu 325 330 335
- Ala Phe His Leu Leu Arg Glu Val Ser Asp Tyr Phe Arg Gln Ser Glu 340 345 350
- Pro His Ser Pro Ile Ser Phe Leu Leu Glu Lys Ala Ile Arg Trp Gly 355 360 365
- Tyr Leu Ser Leu Pro Glu Leu Leu Arg Glu Met Met Ser Glu Gln Asn 370 380
- Gly Asp Ala Leu Ser Thr Ile Phe Asn Ala Ala Gly Leu Asn His Leu 385 390 395 400

Asp Gln Val Leu Leu Pro Glu Val Ser Thr Pro Thr Val Gly Ile Glu 405 410 415

Ser Pro Gln Thr Pro Gln Ala Lys Pro Ser Val Ser Asp Pro Arg Ser 420 425 430

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Asp Ile Gly Asn Gly Thr Asn Ala Asp Ser Gly Met Val Gly Val Ser 50 55 60

Glu Val Ser Val Thr Lys Glu Val Asp Gly Ala Ser Glu Asp Leu Leu 65 70 75 80

Ser Tyr Leu Phe Asn Pro Gly Lys Asp Gly Lys Thr Val Glu Val Ala 85 90 95

Phe Thr Lys Pro Ser Asn Asp Gly Gln Gly Ala Asp Val Tyr Phe Gln 100 105 110

Val Lys Leu Glu Lys Ala Arg Leu Val Ser Tyr Asn Val Ser Gly Thr 115 120 125

Asp Gly Ser Gln Pro Tyr Glu Ser Leu Ser Leu Ser Tyr Thr Ser Ile 130 135 140

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Ser Glu Lys Val Asp Leu Glu Glu Arg Glu Phe Thr Gly Ile Asp Lys
Asp Asn Phe Asp Thr Val Met Gly Gln Ile His Pro Arg Leu Ser Tyr
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Xaa Ile Glu Pro Leu
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